CRF Errors Corrected by the STIC Systems Branch

2/13 OIPE

Serial Number: 09/949947A   **ENTERED**

CRF Processing Date: 2/19/2002
Edited by: _____
Verified by: _____ (STIC staff)

☐ Changed a file from non-ASCII to ASCII

☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐ Edited a format error in the Current Application Data section, specifically:
_____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.

☐ Added the mandatory heading and subheadings for "Current Application Data".

☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
_____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
_____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
_____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included:
_____

☐ Deleted extra, invalid, headings used by an applicant, specifically:
_____

☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as_____.

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically:
_____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically:
_____

☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other:
_____
_____
_____

*Examiner: The above corrections must be communicated to th applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

```
   1 <110> APPLICANT: Brzostowicz, Patricia C.
   2        Cheng, Qiong
   3        DiCosimo, Deana J.
   4        Koffas, Mattheos
   5        Miller, Edward S. Jr.
   6        Odom, J. Martin
   7        Picataggio, Steve
   8        Rouviere, Pierre E.
   9 <120> TITLE OF INVENTION: CAROTENOID PRODUCTION FROM A SINGLE CARBON SOURCE
  10 <130> FILE REFERENCE: CL1903 US NA
C-> 11 <140> CURRENT APPLICATION NUMBER: US/09/941,947A
  12 <141> CURRENT FILING DATE: 2001-09-01
  13 <150> PRIOR APPLICATION NUMBER: 60/229,907
  14 <151> PRIOR FILING DATE: 2000-09-01
  15 <150> PRIOR APPLICATION NUMBER: 60/229,858
  16 <151> PRIOR FILING DATE: 2000-09-01
  17 <160> NUMBER OF SEQ ID NOS: 60
  18 <170> SOFTWARE: Microsoft Office 97
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  22 <212> TYPE: DNA
  23 <213> ORGANISM: Methylomonas 16a
  24 <400> SEQUENCE: 1
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  27        ggcggcttgg cgccttgttt gaattccgca atcggtagtt tgatcgaacg ttataccgaa     180
  28        atcgatccta gcatagaaat catttgctat cgcggcggtt ataaaggcct gttgctgggc     240
  29        gattcttatc cagtaacggc cgaagtgcgt aaaaaggcgg gtgttctgca acgttttggc     300
  30        ggttctgtga tcggcaacag ccgcgtcaaa ttgaccaatg tcaaagactg cgtgaaacgc     360
  31        ggtttggtca agagggtga agatccgcaa aaagtcgcgg ctgatcaatt ggttaaggat     420
  32        ggtgtcgata ttctgcacac catcggcggc gatgatacca atacggcagc agcggatttg     480
  33        gcagcattcc tggccagaaa taattacgga ctgaccgtca ttggtttacc taaaaccgtc     540
  34        gataacgacg tatttccgat caagcaatca ctaggtgctt ggactgccgc cgagcaaggc     600
  35        gcgcgttatt tcatgaacgt ggtggccgaa aacaacgcca acccacgcat gctgatcgta     660
  36        cacgaagtga tgggccgtaa ctgcggctgg ctgaccgctg caaccgcgca ggaatatcgc     720
  37        aaattactgg accgtgccga gtggttgccg gaattgggtt tgactcgtga atcttatgaa     780
  38        gtgcacgcgg tattcgttcc ggaaatggcg atcgacctgg aagccgaagc caagcgcctg     840
  39        cgcgaagtga tggacaaagt cgattgcgtc aacatcttcg tttccgaagg tgccggcgtc     900
  40        gaagctatcg tcgcggaaat gcaggccaaa ggccaggaag tgccgcgcga tgcgttcggc     960
  41        cacatcaaac tggatgcggt caaccctggt aaatggttcg gcgagcaatt cgcgcagatg    1020
  42        ataggcgcgg aaaaaaccct ggtacaaaaa tcgggatact cgcccgtgc ttctgcttcc    1080
  43        aacgttgacg acatgcgttt gatcaaatcg tgcgccgact ggcggtcga gtgcgcgttc    1140
  44        cgccgcgagt ctggcgtgat cggtcacgac gaagacaacg gcaacgtgtt gcgtgcgatc    1200
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50 <212> TYPE: PRT
51 <213> ORGANISM: Methylomonas 16a
52 <400> SEQUENCE: 2
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56                      20                  25                  30
57      Lys Lys Val Ala Ile Leu Thr Ala Gly Gly Leu Ala Pro Cys Leu Asn
58                  35                  40                  45
59      Ser Ala Ile Gly Ser Leu Ile Glu Arg Tyr Thr Glu Ile Asp Pro Ser
60          50                  55                  60
61      Ile Glu Ile Ile Cys Tyr Arg Gly Gly Tyr Lys Gly Leu Leu Leu Gly
62      65                  70                  75                  80
63      Asp Ser Tyr Pro Val Thr Ala Glu Val Arg Lys Lys Ala Gly Val Leu
64                      85                  90                  95
65      Gln Arg Phe Gly Gly Ser Val Ile Gly Asn Ser Arg Val Lys Leu Thr
66                      100                 105                 110
67      Asn Val Lys Asp Cys Val Lys Arg Gly Leu Val Lys Glu Gly Glu Asp
68                  115                 120                 125
69      Pro Gln Lys Val Ala Ala Asp Gln Leu Val Lys Asp Gly Val Asp Ile
70          130                 135                 140
71      Leu His Thr Ile Gly Gly Asp Asp Thr Asn Thr Ala Ala Ala Asp Leu
72      145                 150                 155                 160
73      Ala Ala Phe Leu Ala Arg Asn Asn Tyr Gly Leu Thr Val Ile Gly Leu
74                      165                 170                 175
75      Pro Lys Thr Val Asp Asn Asp Val Phe Pro Ile Lys Gln Ser Leu Gly
76                      180                 185                 190
77      Ala Trp Thr Ala Ala Glu Gln Gly Ala Arg Tyr Phe Met Asn Val Val
78                  195                 200                 205
79      Ala Glu Asn Asn Ala Asn Pro Arg Met Leu Ile Val His Glu Val Met
80          210                 215                 220
81      Gly Arg Asn Cys Gly Trp Leu Thr Ala Ala Thr Ala Gln Glu Tyr Arg
82      225                 230                 235                 240
83      Lys Leu Leu Asp Arg Ala Glu Trp Leu Pro Glu Leu Gly Leu Thr Arg
84                      245                 250                 255
85      Glu Ser Tyr Glu Val His Ala Val Phe Val Pro Glu Met Ala Ile Asp
86                      260                 265                 270
87      Leu Glu Ala Glu Ala Lys Arg Leu Arg Glu Val Met Asp Lys Val Asp
88                  275                 280                 285
89      Cys Val Asn Ile Phe Val Ser Glu Gly Ala Gly Val Glu Ala Ile Val
90          290                 295                 300
91      Ala Glu Met Gln Ala Lys Gly Gln Glu Val Pro Arg Asp Ala Phe Gly
92      305                 310                 315                 320
93      His Ile Lys Leu Asp Ala Val Asn Pro Gly Lys Trp Phe Gly Glu Gln
94                      325                 330                 335
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 95      Phe Ala Gln Met Ile Gly Ala Glu Lys Thr Leu Val Gln Lys Ser Gly
 96                  340                 345                 350
 97      Tyr Phe Ala Arg Ala Ser Ala Ser Asn Val Asp Asp Met Arg Leu Ile
 98              355                 360                 365
 99      Lys Ser Cys Ala Asp Leu Ala Val Glu Cys Ala Phe Arg Arg Glu Ser
100           370                 375                 380
101      Gly Val Ile Gly His Asp Glu Asp Asn Gly Asn Val Leu Arg Ala Ile
102      385                 390                 395                 400
103      Glu Phe Pro Arg Ile Lys Gly Gly Lys Pro Phe Asn Ile Asp Thr Asp
104                  405                 410                 415
105      Trp Phe Asn Ser Met Leu Ser Glu Ile Gly Gln Pro Lys Gly Gly Lys
106                  420                 425                 430
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108                  435
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111 <211> LENGTH: 636
112 <212> TYPE: DNA
113 <213> ORGANISM: Methylomonas 16a
114 <400> SEQUENCE: 3
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116      atggtcatca atcatctgga acatgccgtc cctctggctc gcgcgctagt cgacggtggc     120
117      ttgaaagttt tggagatcac attgcgcacg ccggtggcac tggaatgtat ccgacgtatc     180
118      aaagccgaag taccggacgc catcgtcggc gcgggcacca tcatcaaccc tcataccttg     240
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121      gtcatgcgtt tattggaaaa aggcatcaat gcgatgaaat ctttccggc tgaagccgcc     420
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124      gtcggcggct cctggatggc gccggccgat ctggtagatg ccgaagactg ggcggaaatc     600
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128 <211> LENGTH: 212
129 <212> TYPE: PRT
130 <213> ORGANISM: Methylomonas 16a
131 <400> SEQUENCE: 4
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133      1                 5                  10                  15
134      Val Met Pro Val Met Val Ile Asn His Leu Glu His Ala Val Pro Leu
135                  20                  25                  30
136      Ala Arg Ala Leu Val Asp Gly Gly Leu Lys Val Leu Glu Ile Thr Leu
137              35                  40                  45
138      Arg Thr Pro Val Ala Leu Glu Cys Ile Arg Arg Ile Lys Ala Glu Val
139          50                  55                  60
140      Pro Asp Ala Ile Val Gly Ala Gly Thr Ile Ile Asn Pro His Thr Leu
141      65                  70                  75                  80
142      Tyr Gln Ala Ile Asp Ala Gly Ala Glu Phe Ile Val Ser Pro Gly Ile
143                  85                  90                  95
144      Thr Glu Asn Leu Leu Asn Glu Ala Leu Ala Ser Gly Val Pro Ile Leu
145                  100                 105                 110
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RAW SEQUENCE LISTING　　　　　　DATE: 02/19/2002
PATENT APPLICATION:　US/09/941,947A　　TIME: 13:19:23

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Output Set: N:\CRF3\02192002\I941947A.raw

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149                130                 135                 140
150        Met Leu Lys Ser Leu Gly Gly Pro Leu Pro Gln Val Thr Phe Cys Pro
151        145                 150                 155                 160
152        Thr Gly Gly Val Asn Pro Lys Asn Ala Pro Glu Tyr Leu Ala Leu Lys
153                        165                 170                 175
154        Asn Val Ala Cys Val Gly Gly Ser Trp Met Ala Pro Ala Asp Leu Val
155                        180                 185                 190
156        Asp Ala Glu Asp Trp Ala Glu Ile Thr Arg Arg Ala Ser Glu Ala Ala
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161  <210> SEQ ID NO: 5
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163  <212> TYPE: DNA
164  <213> ORGANISM: Methylomonas 16a
165  <400> SEQUENCE: 5
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167        gcgctgtcca aggaccagct ccagcaactg gctgacgagg tgcgcggcta tctgacccac       120
168        acggtcagca tttccggcgg ccattttgcg gccggcctcg gcaccgtgga actgaccgtg       180
169        gccttgcatt atgtgttcaa taccccgtc gatcagttgg tctgggacgt gggccatcag       240
170        gcctatccgc acaagattct gaccggtcgc aaggagcgca tgccgaccat tcgcaccctg       300
171        ggcggggtgt cagcctttcc ggcgcgggac gagagcgaat acgatgcctt cggcgtcggc       360
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176        agcaagtttt attcgtcggt gcgggaagag agcaagaaag ctctggccaa gatgccgtcg       660
177        gtgtgggaac tggcgcgcaa gaccgaggaa cacgtgaagg gcatgatcgt gcccggtacc       720
178        ttgttcgagg aattgggctt caattatttc ggcccgatcg acggccatga tgtcgagatg       780
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182        tataccgagg tgttcggccg ctggctgtgc gacatggcgg ctcaagacga gcgcttgctg      1020
183        ggcatcacgc cggcgatgcg cgaaggctct ggtttggtgg aattctcaca gaaatttccg      1080
184        aatcgctatt tcgatgtcgc catcgccgag cagcatgcgg tgaccttggc cgccggccag      1140
185        gcctgccagg gcgccaagcc ggtggtggcg atttattcca ccttcctgca acgcggttac      1200
186        gatcagttga tccacgacgt ggccttgcag aacttagata tgctctttgc actggatcgt      1260
187        gccggcttgg tcggcccgga tggaccgacc catgctggcg cctttgatta cagctacatg      1320
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191        caccacggca gccgcatcgc cattctggcc tggggcagca tggtcacgcc tgccgtcgaa      1560
192        gccggcaagc agctgggcgc gacggtggtg aacatgcgtt tcgtcaagcc gttcgatcaa      1620
193        gccttggtgc tggaattggc caggacgcac gatgtgttcg tcaccgtcga ggaaaacgtc      1680
194        atcgccggcg cgctggcag tgcgatcaac accttcctgc aggcgcagaa ggtgctgatg      1740
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OIPE

RAW SEQUENCE LISTING   DATE: 02/10/2002
PATENT APPLICATION: US/09/941,947A   TIME: 13:19:55
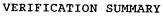
Input Set : A:\CL1903 US NA revised seq list .txt
Output Set: N:\CRF3\02102002\I941947A.raw

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 3 <110> APPLICANT: Brzostowicz, Patricia C.
 4        Cheng, Qiong
 5        DiCosimo, Deana J.
 6        Koffas, Mattheos
 7        Miller, Edward S. Jr.
 8        Odom, J. Martin
 9        Picataggio, Steve
10        Rouviere, Pierre E.
14 <120> TITLE OF INVENTION: CAROTENOID PRODUCTION FROM A SINGLE CARBON SOURCE
18 <130> FILE REFERENCE: CL1903 US NA
21 <140> CURRENT APPLICATION NUMBER: US/09/941,947A
21 <141> CURRENT FILING DATE: 2001-09-01
21 <150> PRIOR APPLICATION NUMBER: 60/229,907
22 <151> PRIOR FILING DATE: 2000-09-01
24 <150> PRIOR APPLICATION NUMBER: 60/229,858
25 <151> PRIOR FILING DATE: 2000-09-01
27 <160> NUMBER OF SEQ ID NOS: 60
29 <170> SOFTWARE: Microsoft Office 97
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Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES

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3085 <210> SEQ ID NO: 60
3086 <211> LENGTH: 19
3087 <212> TYPE: DNA
3088 <213> ORGANISM: Artificial Sequence
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3091 <223> OTHER INFORMATION: primer
3093 <400> SEQUENCE: 60
3094 tagctcgagt cacgcttgc                                          19
E--> 3097 (52) delete
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VERIFICATION SUMMARY   DATE: 02/10/2002
PATENT APPLICATION: US/09/941,947A   TIME: 13:19:56

Input Set : A:\CL1903 US NA revised seq list .txt
Output Set: N:\CRF3\02102002\I941947A.raw

L:21 M:270 C: Current Application Number differs, Replaced Current Application No
L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:3097 M:254 E: No. of Bases conflict, LENGTH:Input:52 Counted:19 SEQ:60

US 0994194708P1

Creation date: 28-08-2003
Indexing Officer: FPLUMMER - FRANCIS PLUMMER
Team: OIPEBackFileIndexing
Dossier: 09941947

Legal Date: 27-09-2002

No.	Doccode	Number of pages
1	IDS	3
2	NPL	3

Total number of pages: 6

Remarks:

Order of re-scan issued on ....